

Fig 1

rGFra-5(A) : **MLSGAYLRVLNERPGQAVLWSLGCQRGSSASSTEGNRCVEAAEACTADEQC** : 50
 rGFra-5(B) : **MLSGAYLRVLNERPGQAVLWSLGCQRGSSASSTEGNRCVEAAEACTADEQC** : 50

rGFra-5(A) : **QQLRSEYVAQCLGRAGWRGPGSCVRSRRCRRALRRFFARGPPALTHALLFC** : 100
 rGFra-5(B) : **QQLRSEYVAQCLGRAGWRGPGSCVRSRRCRRALRRFFARGPPALTHALLFC** : 100

rGFra-5(A) : **GCEGPACAERRRQTFAPACAFSGPQLAPPSCCLKPLDRCSRRRCRPRLEA** : 150
 rGFra-5(B) : **GCEGPACAERRRQTFAPACAFSGPQLAPPSCCLKPLDRCSRRRCRPRLEA** : 150

rGFra-5(A) : **FQASCAPAPGSRDGCPEEGGPRCLRAYAGLVGTVVTNLYLDNV SARVAPW** : 200
 rGFra-5(B) : **FQASCAPAPGSRDGCPEEGGPRCLRAYAGLVGTVVTNLYLDNV SARVAPW** : 200

rGFra-5(A) : **CGCEASGNRRREECEAFRKLFTTRNPCLDGAIQAFDSSQPSVLQDQWNP YQN** : 250
 rGFra-5(B) : **CGCEASGNRRREECEAFRKLFTTRNPCLDGAIQAFDSSQPSVLQDQWNP YQN** : 250

rGFra-5(A) : **AGCCFLNVSSMSILTALALQALL** : 273
 rGFra-5(B) : **AGQAKVEA-----** : 258

Fig 2

CGFA-5(A)	-----MISCHLVINERPQ-----	16
CGFA-5(B)	-----MISCHLVINERPQ-----	16
CGFA-4	-----MISCHLVINERPQ-----	57
CGFA-3	-----MISCHLVINERPQ-----	120
CGFA-2	-----MISCHLVINERPQ-----	69
CGFA-1	-----MISCHLVINERPQ-----	63

[illegible][illegible][illegible]

CGFLA5 (A) :	CGFLAYSSMSLTALNALQLL-----	273
CGFLA5 (B) :	CGFLA5 (B) :	258
CGFLA5 (B) :	CGFLA5 (B) :	431
CGFLA-4 :	CGFLA-4 :	397
CGFLA-3 :	CGFLA-3 :	444
CGFLA-2 :	CGFLA-2 :	468
CGFLA-1 :	CGFLA-1 :	

File 3

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Fig 4

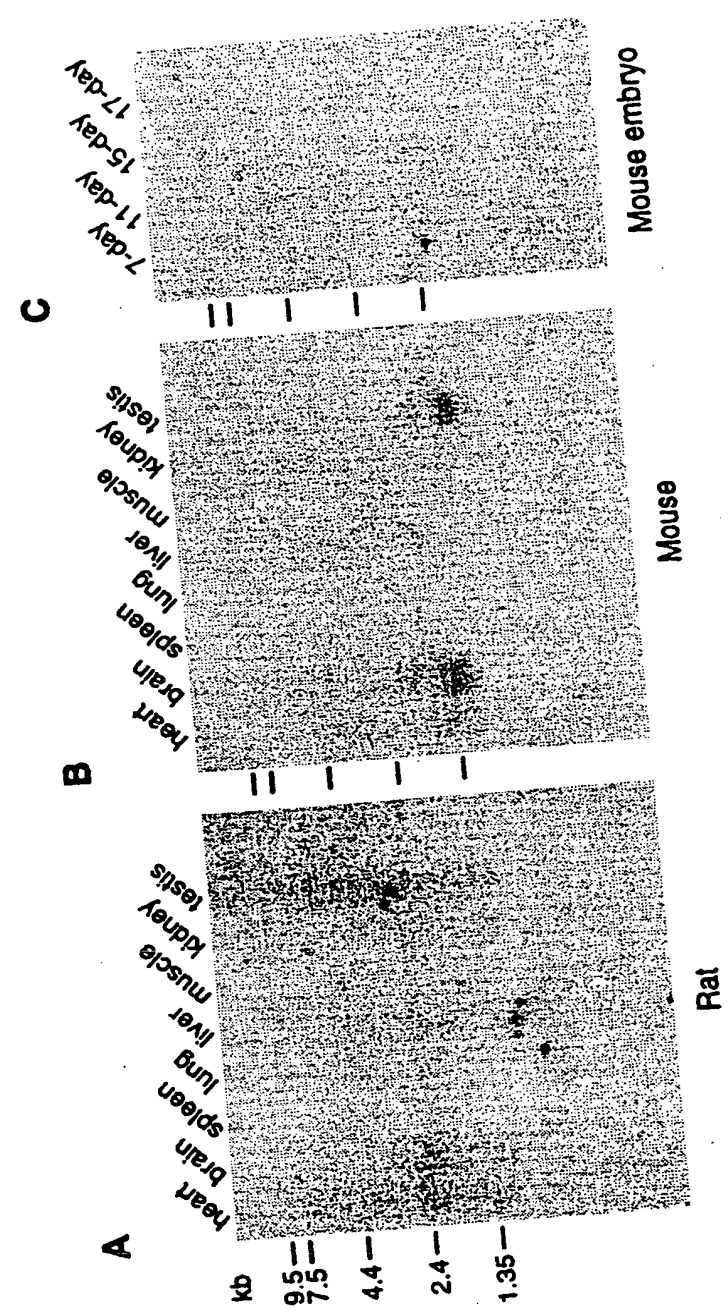
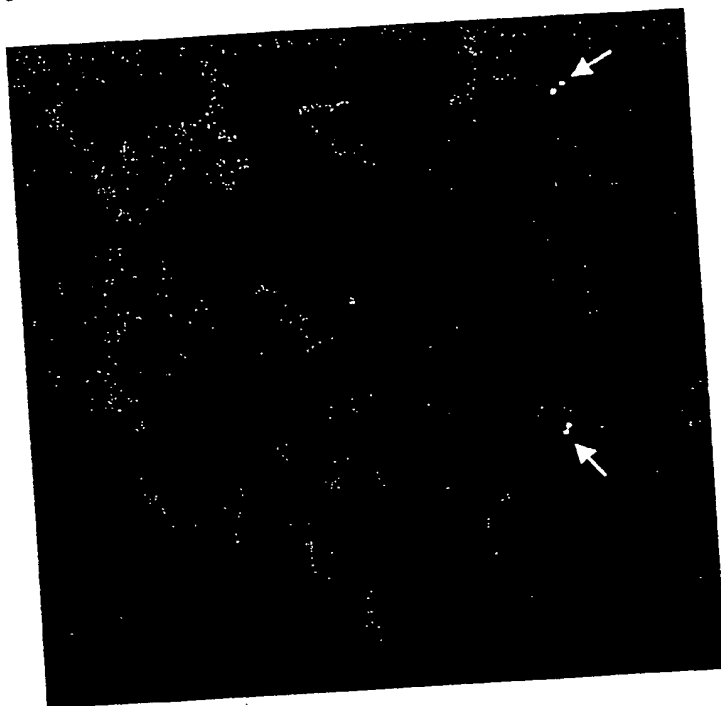


Fig 5**A****B**